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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*

1: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Sequence 1, Appli Sequence 296, App	Sequence 17684, A	565	Sequence 2, Appli	24,	Sequence 33, Appl	285		Sequence 102915,	Sequence 1, Appli	Sequence 120, App	Sequence 120, App	Sequence 8174, Ap	Sequence 8559, Ap	Sequence 4589, Ap	58879	Sequence 600, App	Sequence 10421, A				Sequence 1436, Ap	10	10,	19	326,	27,		Sequence 240, App	Sequence 100851,	Sequence 68, Appl	Sequence 879, App	GENERAL INFORMATI

ALIGNMENTS

US-10-617-962-2

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Sequence 2, Application US/10617962
Publication No. US2004005536A1
GENERAL INFORMATION:
APPLICANT: Commonwealth Scientific and Industrial Research Organisation
APPLICANT: EAST, Peter David
TITLE OF INVENTION: Toxin Genes from the Bacteria Xenorhabdus nematophilus and Photo:
TITLE OF INVENTION: Tuminescens
FILE REFERENCE: 050179-0076
CURRENT FILING DATE: 2003-07-14
PRIOR APPLICATION NUMBER: US/10/617,962
CURRENT FILING DATE: 2003-07-14
PRIOR APPLICATION NUMBER: US/09/463,048A
PRIOR FILING DATE: 1998-07-17
PRIOR APPLICATION NUMBER: PCT/AU98/00562
PRIOR FILING DATE: 1998-07-17
NUMBER OF SEQ ID NOS: 20
SOFTWARE: Patentin version 3.1
SEQ ID NO 2
LENGTH: 1008
TYPE: DNA
ORGANISM: Photorhabdus luminescens
US-10-617-962-2
Query Match
Best Local Similarity 100.0%; Score 1008; DB 17; Length 1008;
Best Local Similarity 100.0%; Pred. No. 1.9e-198;
Matches 1008; Conservative 0; Mismatches 0; Gaps 0;
Matches 1008; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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RESULT 2
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; Publication No. US20040055036A1
; GENERAL INFORMATION:
; APPLICANT: COmmonwealth Scientific and
; APPLICANT: EAST, Peter David
; TITLE OF INVENTION: Toxin Genes from th
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CURRENT APPLICATION NUMBER: US/10/617,962

CURRENT FILING DATE: 2003-07-14

PRIOR APPLICATION NUMBER: US/09/463,048A

PRIOR FILING DATE: 2002-12-13

PRIOR APPLICATION NUMBER: PO 8098

PRIOR FILING DATE: 1998-07-17

PRIOR FILING DATE: 1997-07-17

NUMBER OF SEQ ID NOS: 20

SOFTWARE: Patentin version 3.1

SEQ ID NO 1

LENCTH: 1107

TYPE: DNA

ORGANISM: Xenorhabdus nematophilus
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Best Local Similarity
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                                                                                                          TTATATTGAGAGAAAAAACTTTCTAAACCTTTCTTTCGACCACCGCAGAGATTACCTGA
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TGGCAGAACAGGCTACTTGGCCGGTCCAACAAAAAGCGCCTAAATTGCCAACAACGTCTTC
                                                                          GGGAAAAAGGATTTTTTCAAAAAATTTTTACAATGGTCTAGAAACTAAATCATTAGAGAA
                                                                                                                                                                                           GGGAAGGAGTTTTTTCAAAAAACTTTTACAATGGATTAGAGGCTAAATCATTAGAGAA
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CURRENT APPLICATION NUMBER: US/10/617,962
CURRENT FILING DATE: 2003-07-14
PRIOR APPLICATION NUMBER: US/09/463,048A
PRIOR FILING DATE: 2002-12-13
PRIOR FILING DATE: 2002-12-13
PRIOR APPLICATION NUMBER: PCT/AU99/00562
PRIOR FILING DATE: 1998-07-17
PRIOR APPLICATION NUMBER: PO 8088
PRIOR FILING DATE: 1997-07-17
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Publication No. US20040055036A1
GENERAL INFORMATION:
APPLICANT: Commonwealth Scientific and Industrial Research Organisation
APPLICANT: EAST, Peter David
                                                                                                                                                                                                                                                                                                                                                                       Best Loca
Matches
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SOFTWARE: PatentIn version
SEQ ID NO 5
LENGTH: 1205
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Toxin Genes from the Bacteria Xenorhabdus nematophilus and TITLE OF INVENTION: luminescens
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Xenorhabdus nematophilus
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                                    ATACATGTCTAATCTTTTATCTGCACTATTGGGTGAGAATGGTGATCAAATTTCTAAAGA
                                                                                                   CTCTCAACTTCCAGAAGACTATTATAAAAATAAGGATAACACAGGTAGAATTTATCAAGA
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                                    ÚS-10-473-126-386
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                                                                                                                                                                                                 Sequence 386, Application US/10473126
Publication No. US20040234973A1
GENERAL INFORMATION:
APPLICANT: Epigenomics AG
TITLE OF INVENTION: Methods and nucleic acids for the analysis of
TITLE OF INVENTION: proliferative disorders
FILE REFERENCE:
                                                                                 CURRENT APPLICATION NUMBER: US/10/473,126
CURRENT FILING DATE: 2003-09-26
NUMBER OF SEQ ID NOS: 1258
SEQ ID NO 386
LENGTH: 8056
TYPE: DNA
ORGANISM: Artificial Sequence
   Query Match
                                                    OTHER INFORMATION:
                                                                     FEATURE:
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RESULT 5
US-10-235-192A-27
J Sequence 27, Application US/10235192A
Publication No. US20040043389A1
GENERAL INFORMATION:
                                          문
                                                                                                                         Query Match
Best Local Similarity
Matches 263; Conserv
                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity Matches 226; Conserv
                                                                                                                                                                                                      FEATURE:
NAME/KEY: misc feature
LOCATION: 145329-145428, 187896-187995,
OTHER INFORMATION: N = any nucleotide
0-10-235-192A-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: McCarthy, Jeanette
TITLE OF INVENTION: Methods and Compositions for Identifying
TITLE OF INVENTION: Risk Factors for Abnormal Lipid Levels and
TITLE OF INVENTION: and Disorders Associated Therewith
FILE REFERENCE: MMI-011
                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/235,192A
CURRENT FILING DATE: 2002-09-04
NUMBER OF SEQ ID NOS: 49
                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                 ATTAACACCTGATGATAGAAGTGGATATCCACCCGTTGAAAAGCAAATAGCAGGAGATAT
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                                                                                                                       Score 50.6; DB 17;
Pred. No. 10;
0; Mismatches 354;
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0; Mismatches
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  US-10-240-453-95
                                                                                                                                                         FILE REFERENCE: 5013.1009

CURRENT APPLICATION NUMBER: US/10/240,453

CURRENT FILING DATE: 2002-10-02

PRIOR APPLICATION NUMBER: PCT/EP01/03973

PRIOR FILING DATE: 2001-04-06

PRIOR APPLICATION NUMBER: DE 10019058.8

PRIOR FILING DATE: 2000-04-06

PRIOR FILING DATE: 2000-04-07

PRIOR APPLICATION NUMBER: DE 10019173.8

PRIOR FILING DATE: 2000-04-07

PRIOR APPLICATION NUMBER: DE 10032529.7

PRIOR APPLICATION NUMBER: DE 10043826.1

PRIOR PILING DATE: 2000-09-01
                                                                                             NUMBER OF SEQ ID NOS: 350
SEQ ID NO 95
LENGTH: 11394
                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: ITTLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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APPLICANT: PIEPE
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                   FEATURE:
OTHER INFORMATION:
                                                           ORGANISM: Artificial Sequence
                                                                                 TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alexander
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                   sapiens)
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PRIOR APPLICATION NUMBER: PCT/US99/01972
PRIOR FILING DATE: 1999-01-29
PRIOR PRIOR POLING DATE: 1998-01-30
PRIOR FILING DATE: 1998-01-30
NUMBER OF SEO ID NOS: 181
SOFTWARE: Patentin version 3.2
SEQ ID NO 73
PEROTH: 1866
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                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 4.8%;
Best Local Similarity 47.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                   Matches 249;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Cadsell, Gail
APPLICANT: Chen, Ellson
APPLICANT: Glass, Jennifer
APPLICANT: Glass, Jennifer
APPLICANT: Hainer, Cheryl
APPLICANT: Hainer, Cheryl
APPLICANT: Lifthowitz, Elliot
TITLE OF INVENTION: NUCLEIC ACID PROBES AND METHOD FOR DETECTING UREAPLASMA
TITLE OF INVENTION: UREALYTICUM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches 120; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/349,680
CURRENT FILING DATE: 2003-01-23
PRIOR APPLICATION NUMBER: US 09/601,198
PRIOR FILING DATE: 2000-12-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: UREALYTICUM FILE REFERENCE: UAB-13403/22
                                            1217
                                                                                                                                                                                                                       1097
                                                                                                                               1157 ATATTACACAAAAATAATTAAAAATCTTTTATTAGATGCTTTCAATAATAATGATGTTC
                                                                                                                                                                                                                                                                                                           1037 ATGAAAAAAACAACAAATAAGAATTATTTTAGATCTTTTGATAAAGAATTGAATTATT 1096
                                                                                                                                                                         241 -- ATTTATCAAGAATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATC
                                                                                                                                                                                                                                                               182 TTCATAATGTATCTAAACTTCCTGATGACTATTATAAGAATAAAGAGACTGCTGAGAGA-
                                                                                                                                                                                                                                                                                                                                                     122 ATGGAATTGAATATCGAGCTAAGAAAATAATTAGCTTACGCTTTGGCTGTAAGTGGTA 181
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  359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             244 TATCAAGAATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAAATT 303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       184 CATAATGTATCTAAACTTCCTGATGACTATTATAAGAATAAAGAGACTGCTGAGAGAATT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TGATATTTÄTTAGTAATGATTGTATTTÄGTGTTTÄGAGTTTTTTGGTTTTTÄÄTÄT 554
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ATCCTCAAAACATTTGGAATGTTCCTGAGCTTGAAAATAAACCATTGAGTGCTTATTCAG
                                            TTCAAAAATATATTGATCATAATTTAGATATTAATATAAAAATTTCAAAGGAAGCAT
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                                                                                     AAATTTCTAAAGATATGGCAAATGGTTTTTATAAGAATGAACTGGATTTTGAAGGTCAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/10349680
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                                                                                                                                                                                                                                                                                                                                                                                               Score 48.8; DB 16; Length 1866; Pred. No. 4.4; 0; Mismatches 267; Indels 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches 117;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 49.8; DE Pred. No. 5.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 15;
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US-10-311-455-597, Application US/10311455
; Sequence 597, Application US/10311455
; Publication No. US20030143606A1
; GENERAL INFORMATION:
APPLICANT: OLEK, Alexander
; APPLICANT: PIEPENBROCK, Christian
APPLICANT: BERLIN, Kurt
; APPLICANT: BERLIN, Kurt
; TITLE OF INVENTION: Diagnosis of Diseases As
; TITLE OF INVENTION: Cytosine methylation
; FILE REFERENCE: 5013.1014
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; LOCATION: (2)...(1507)
US-10-301-533-23
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US-10-301-533-23
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APPLICANT: Reed, John C.
TITLE OF INVENTION: Microbial Sumo Protease Ho
FILE REFERENCE: P-LJ 5488
CURRENT APPLICATION NUMBER: US/10/301,533
CURRENT FILING DATE: 2002-11-20
PRIOR APPLICATION NUMBER: US 60/331,895
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 51
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
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LENGTH: 2767
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Plasmodium
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1277 AATTAATAGAATTAGTAATTATTTTAAAAACCAAAAAAACCCAAGAGAGATTCTAATGAAT 1336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1455 AAATCCTATTACTAAAGAGATGTTAGGACCAAGTGAAAAATATTTAAAAAAGGTTAAAAAA
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                                                                                                                                                                                                                                                                  682 TGATGAAGTATACAATAATATAGATGATGACTATAAA 718
                                                                                                                                                                                                                                                                                                      363 TCAAAACATTTGGAATGTTCCTGAGCTTGAAAATAAA 399
                                                                                                                                                                                                                                                                                                                                            243 TTATCAAGAATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     594 GGCTAAATCATTAGAGAATTATTATTGAGAGAAAAAAACTTTCTAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    , 68
APPLICATION NUMBER: US/10/311,455
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Pred. No. 6.7;
0; Mismatches
                                                        Associated with the
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RESULT 10
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LENGTH: 16217
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                         Sequence 531, Application US/08781986A Publication No. US20030054436A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                     GENERAL INFORMATION:
APPLICANT: Charles Kunsch
TITLE OF INVENTION: Staphylococcus
NUMBER OF SEQUENCES: 5255
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT FILING DATE: 2002-12-16
PRIOR APPLICATION NUMBER: PCT/BP01/07537
PRIOR FILING DATE: 2001-07-02
PRIOR APPLICATION NUMBER: DE 10032529.7
PRIOR FILING DATE: 2000-06-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: DE 10043826.1 PRIOR FILING DATE: 2000-09-01
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OTHER INFORMATION: n is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
OTHER INFORMATION:
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STREET:
CITY: I
                             ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          124 GGAATTGAATATCGAGCTAAGAAAATAATATTAGCTTACGCTTTGGCTGTAAGTGGTATT 183
 Rockville
                                                                                                                                                                                                                      AGTGGGAATTTTATAGAGAATATTAAATTT 3004
                                                                                                                                                                                                                                           TTAGAGAATTATTTGAGAGAAAAAAACTT
                                                                                                                                                                                                                                                                                 AGTAGTAGTTTATTTTTTGATTTTAGTTAAAAAGTGTATAAAATATGATATATGGAATTT 2974
                                                                                                                                                                                                                                                                                                                                             CAATCAAATGCCGCAAGATTTTTTAAATTAATTGATTTCTTATTTACCTTATCTGCTGTA 543
                                                                                                                                                                                                                                                                                                                                                                                                                                          GATAAATTATTAGCACTATATTTTTTCTCTGTACAGGAAATTCCACTGGAGGAAAATCAA 483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAGGTAATTAATATTTTGGTTATTGAATTTTÄTATTATÄTAAATTTATTAATAAATTTAAT 2794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CAAAACATTTGGAATGTTCCTGAGCTTGAAAATAAACCATTGAGTGCTTATTCAGATGAC 423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TATCAAGAATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAAATT 303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAAATAATGTATTTTATTAAAAAATTAAAATTAGAAAATGTAATATAAAGATGTAATG 2554
                                                                                                                                                                                                                                                                                                            ACTTCACTGGGAAGGAGGATTTTTTCAAAAAACTTTTACAATGGATTAGAGGCTAAATCA
                                                                                                                                                                                                                                                                                                                                                                                                            AGTATTTAGTTTAGATTTTATTATTATTTTGTTAATTTAGTTTAGTTATAGAGAAATTA 2854
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TCTAAAGATATGGCAAATGGTTTTTATAAGAATGAACTGGATTTTGAAGGTCAATATCCT 363
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9410 Key
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              Genome Sciences,
West Avenue
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Pred. No. 17;
0; Mismatches 289;
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COUNTRY:
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RESULT 11
US-10-39-624-531
JS-10-39-624-531
JS-20-39-624-531
JS-20-39-624-531
JS-20-64-631
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (301) 309-85
INFORMATION FOR SEQ ID NO:
Gil H. Choi

Patrick S. Dillon

Craig A. Rosen

Craig A. Rosen

Steven C. Barash

Michael R. Fannon

TITLE OF INVENTION: Staphylococcus auret

NUMBER OF SEQUENCES: 5256

CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences, Ir

STREET: 9410 Key West Avenue

CITY: Rockville

STATE: Marvland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 942 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 20850
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Benson, Bob
REGISTRATION NUMBER: 30,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
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Pred. No. 6.6;
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RESULT 12
US-10-211-179-11
                                                        GENERAL INFORMATION:

APPLICANT: Nicholas M. Dean

APPLICANT: Kenneth W. Dobie

TITLE OF INVENTION: ANTISENSE MODULATION OF PHOSPHOTYROSYL PHOSPHATASE ACTIVATOR EXPR

FILE REFERENCE: PTS-0011

CURRENT APPLICATION UMBER: US/10/211,179

CURRENT FILING DATE: 2002-08-01

NUMBER OF SEQ ID NOS: 119

OTHER INFORMATION: n = a, t, c, or 9

US-10-211-179-11
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  Query Match
Best Local Similarity
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SEQUENCE CHARACTERISTICS:
LENGTH: 942 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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APPLICATION NUMBER: 08/781,986
FILING DATE: January 3, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Mark J. Hyman
REGISTRATION NUMBER: 46,789
REGERENCE/DOCKET NUMBER: PB248P1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/956,171
FILING DATE: 27-Dec-2002
PRIOR APPLICATION NUMBER: 08/956,171
FILING DATE: October 20, 1997
APPLICATION NUMBER: 60/009,861
APPLICATION NUMBER: 5, 1996
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MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS Version 6.2
SOFTWARE: ASCII Text
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APPLICATION NUMBER: US/10/329,624
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TELEFAX: (301) 309-8439
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.78;
4.7%;
42.8%;
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Pred. No. 6.6;
0; Mismatches 151;
  Score 47.4;
Pred. No. 44;
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                    DB 17;
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                  Length 158001;
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                                                                                                                                                                                                                                                                                                                AAAATGATACTGACAATACAATGACGGTTCATAGTGTTGGAACTCATTATAAAAATA
                                             GTAACGCTCCACAGACAGCAGCGAAAAGTATTGGTACAGCCAGTGGCAGTAACCTGGAAA
                                                                                                                                          CTAATATGGAAGGGGCTGCAAAACAAAAGTATAGTTCATTTATAAAAGAGGTACAAAAGG
                                                                                                                                                                                          CGCCTAAATGGAGAGTGAGTTTTAAAGAACTTAAAAAATAACAAATCTAGGAATGGATTTT
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                                                                  AATTGCCGAATAATTTATATAGTGTGAGGCTAAGCCAAAAAGACAGGGTAACCTTTACTC
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RESULT 13
US-09-991-936-879/c
US-09-991-936-879/, Application US/09991936
; Publication No. US20030073827A1
; GENERAL INFORMATION:
; APPLICANT: Brandt, Kevin S.

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RESULT 14
US-10-257-166-68
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; ORGANISM: Ctenocephalides felis
US-09-991-936-879
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                                                                                                                                                                                                                                                                                  Sequence 68, Application US/10257166
Publication No. US20040023230A1
GENERAL INFORMATION:
APPLICANT: OLEK, Alexander
APPLICANT: PIEPEMBROCK, Christian
APPLICANT: BERLIN, Kurt
TITLE OF INVENTION: Method and Nucleic Acids for Analysing the Methylation
TITLE OF INVENTION: Genes Implicated in Pharmacogenomics
FILE REFERENCE: 5013.1011
CURRENT APPLICATION NUMBER: US/10/257,166
CURRENT FILING DATE: 2002-10-07
CURRENT FILING DATE: 2002-10-07
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Best Local Similarity
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APPLICANT: Stinchcomb, Dan T.
APPLICANT: Wisnewski, Nancy
APPLICANT: Wisnewski, Nancy
TITLE OF INVENTION: FLEA HEAD, NERVE CORD, HINDGUT AND MALPIGHIAN TUBULE
TITLE OF INVENTION: NUCLEIC ACID MOLECULES, PROTEINS AND USES THEREOF
FILE REFERENCE: FC-6-C1
CURRENT APPLICATION NUMBER: US/09/991,936
CURRENT FILING DATE: 2001-11-21
PRIOR APPLICATION NUMBER: US/09/543,668
PRIOR EILING DATE: 2000-04-07
PRIOR FILING DATE: 2000-04-07
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PRIOR FILING DATE: 1999-04-09
NUMBER OF SEQ ID NOS: 1959
                                                                                                                                2000-09-01
NUMBER OF SEQ ID
                                                                                                                                                                                                  PRIOR FILING DATE:
                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: PCT/EP01/07470
OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
                                           TYPE: DNA ORGANISM: Artificial Sequence
                         FEATURE:
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                                                                                    ENGTH: 11052
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DE 10043826.1
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49.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 100851
LENGTH: 260
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APPLICANT:
APPLICANT:
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APPLICANT: Kovalic David
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                  FEATURE:
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                                                              CGTTTGGATTTTTTAAAAAAATGAGCATAAACAAAAGTTTGGTCATCTTAAGGGGGGAAA
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Copyright (c) 1993 - 2005 Compugen Ltd.
US-09-463-048A-2
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US-08-569-168-3
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US-09-806-708B-23
US-09-956-171E-531
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301 АТТТСТАЛАСАТАТССАЛАТСТТТТТАТЛАСЛАЛСТАЛССТТТССАТТТССА	241 ATTTATCAAGAATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAA 	121 TATGGAATTGAATATCGAGCTAAGAAAATAATATTAGCTTACGCTTTGGCTGTAAGTGGT	1 ATGGTTATACAATTAACACCTGATGATAGAAGTGATATCACCCCGTTGAAAAGCAAATA	Query Match 100.0%; Score 1008; DB Best Local Similarity 100.0%; Pred. No. 6.7e- Matches 1008; Conservative 0; Mismatches	SULT 1 -09-463-048A-2 Sequence 2, Application US/09463048A Patent No. 6630619 GENERAL INFORMATION: APPLICANT: Commonwealth Scientific and Industrial APPLICANT: EAST, Peter David TITLE OF INVENTION: Toxin Genes from the Bacteria TITLE OF INVENTION: 1uminescens FILE REFERENCE: 050179-0076 CURRENT FILING DATE: 2002-12-13 PRIOR APPLICATION NUMBER: US/09/463,048A CURRENT FILING DATE: 1998-07-17 PRIOR APPLICATION NUMBER: PO 8088 PRIOR FILING DATE: 1998-07-17 RIOR FILING DATE: 1997-07-17 NUMBER OF SEQ ID NOS: 20 SOFTWARE: Patentin version 3.1 SEQ ID NO 2 CHENGTH: 1008 TYPE: DNA ORGANISM: Photorhabdus luminescens -09-463-048A-2	ALIGNMENTS	8 41.8 4.1 9 41.8 4.1 1 0 41.8 4.1 1 1.4 4.1 2 41.4 4.1 4 4.1 4 4.1 4 4.1 4 4.1 4 4.1 4 4.1 4 4.1 4 4.1	2 42.6 4.2 540 4 4.2 6 4.2 627 4 4 42.4 4.2 627 4 4 42.4 4.2 228896 4 5 42.2 4.2 228896 4 7 42.2 4.2 5136 4	8 43.8 4.3 665 2 US-08-883-795A 9 43.2 4.3 1470 4 US-09-830-217- 0 43.2 4.3 1470 4 US-10-278-946- 1 42.8 4.2 7425 3 US-09-453-702B
GAACTGGATTTTGAAGGTCAATAT 360	CTATTAGGTGAAAATGGTGATCAA 300 	GCTTACGCTTTGGCTGTAAGTGGT 180	GATATCCACCGTTGAAAAGCAAATA 60	3 4; Length 1008; 244; 0; Indels 0; Gaps 0;	tal Research Organisation ria Xenorhabdus nematophilus and		Sequence Sequence Sequence Sequence Sequence Sequence Sequence	Sequence 2 Sequence 1 Sequence 1 Sequence 2 Sequence 3 Sequence 3	Sequence Sequence Sequence Sequence

Photo

301 ATTTCTAAAGATATGGCAAATGGTTTTTATAAGAATGAACTGGATTTTTGAAGGTCAATAT 360

Query Match 65.9%; Score 663.8; DB 4; Length 1107; Best Local Similarity 79.4%; Pred. No. 2.6e-157;	; LENGTH: 1107 ; TYPE: DNA ; ORGANISM: Xenorhabdus nematophilus US-09-463-048A-1	; PRIOR FILING DATE: 1997-07-17 ; NUMBER OF SEQ ID NOS: 20 ; SOFTWARE: PatentIn version 3.1 ; SEQ ID NO 1	AFE AF	; TITLE OF INVENTION: Toxin Genes from the Bacteria Xenorhabdus nematophilus and Photo; TITLE OF INVENTION: luminescens ; FILE REFERENCE: 050179-0076 ; FILE REFERENCE: 050179-0076 ; CÜRRENT APPLICATION NUMBER: US/09/463,048A	; Patent No. 6630619 ; GENERAL INFORMATION: ; APPLICANT: Commonwealth Scientific and Industrial Research Organisation ; APPLICANT: EAST, Peter David	RESULT 2 US-09-463-048A-1 ; Sequence 1, Application US/09463048A	Qy 961 AATACAATGACGGTTCATAGTGTTGGAACTCATTATAAAATATATGA 1008 	QY 901 TTATATAGTGTGAGGCTAAGCCAAAAAGACAGGGTAACCTTTACTCAAAATGATACTGAC 960	Qy 841 ACAGCAGCGAAAAGTATTGGTACAGCCAGTGGCAGTAACCTGGAAAAATTGCCGAATAAT 900	Qy 781 GCTGCAAAACAAAAGTATAGTTCATTTATAAAAGAGGTACAAAAGGGTAACGCTCCACAG 840	Qy 721 GTGAGTTTTAAAGAACTTAAAAATAACAAATCTAGGAATGGATTTTCTAATATGGAAGGG 780	Qy 661 AGATTACCTGATGGCAGAATAGGTTATTTGGCTGGACCAACAGAAGCGCCTAAATGGAGA 720	Qy 601 TCATTAGAGAATTATATTGAGAGAAAAAAACTTTCTAAACCTTTCTTT	OY 541 GTAACTTCACTGGGAAGGAGGATTTTTTCAAAAAACTTTTACAATGGATTAGAGGCTAAA 600	Oy 481 CAACAATCAAATGCCGCAAGATTTTTTAAATTAATTTATTT	Qy 421 GACGATAAATTATTAGCACTATATTTTTTTCTCTGTACAGGAAATTCCACTGGAGGAAAAT 480	Qy 361 CCTCAAAACATTIGGAATGITCCIGAGCTIGAAAATAAACCATTGAGTGCTTATTCAGAT 420	Db 301 ATTTCTAAAGATATGGCAAATGGTTTTTATAAGAATGAACTGGATTTTGAAGGTCAATAT 360
Db 1077 AAGTGTTGG	918 1017	Qy 861 TACAGCCAG Db 957 TACAAAAAA	Oy 801 TTCATTTAT	2 Qy 741 Db 837	Qy 715Db 777 TACAGCAAC	Qy 672 TGGCAGAAT	Qy 612 TTATATTGA Db 657 TTATATTGA	Qy 552 GGGAAGGAG Db 597 GGGAAAAAAG	Oy 492 TGCCGCAAG Db 537 TGCAGCAAA	Oy 432 ATTAGCACT	Oy 372 TTGGAATGT	Qy 312 TATGGCAAA Db 357 TATGGCAAA	Qy 252 ATATATGTC	Qy 192 ATCTADACT	Qy 132 ATATCGAGG	Qy 72 AGTACGTAT 	Qy 12 ATTAACACC Db 57 AGTAACGCC	Matches 835; Conservative
I I I I I I I I I I	AAGCCAAAAACACAGGGTAACCTTTACTCAAAATGATACTGACAATACAATGACGGTTCA 977	TACAGCCAGTGGCAGTAACCTGGAAAATTGCCGAATAATTTATATAGTGTGAGGCT 917		AAATAACAAATCTAGGAATGGATTTTCTAATATGGAAGGGGCTGCAAAACAAAAGTATAG 800 	TACAGCAACAACGTCTACAGCAGCTTCATCTAATTGGAGAGTTAGTT	TGGCAGAATAGGTTATTTGGCTGGACCAACAGAAGCGCCTAAA714 	TTATATTGAGAGAAAAAACTTTCTAAACCTTTCTTTCGACCACCGCAGAGATTACCTGA 671	GGGAAGGAGGATTTTTTCAAAAAACTTTTACAATGGATTAGAGGGCTAAATCATTAGAGAA 611 	TGCCGCAAGATTTTTTAAATTAATTGATTTCTTATTTACCTTATCTGCTGTAACTTCACT 551	ATTAGCACTATATTTTTTCTCTGTACAGGAAATTCCACTGGAGGAAAATCAACAATCAAA 491 	TTGGAATGTTCCTGAGCTTGAAAATAAACCATTGAGTGCTTATTCAGATGACGATAAATT 431 	TATGGCAAATGGTTTTTATAAGAATGAACTGGATTTTGAAGGTCAATATCCTCAAAACAT 371 	ATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAAATTTCTAAAGA 311	192 ATCTAAACTTCCTGATGACTATTATAAGAATAAAGAGACTGCTGAGAGAATTTATCAAGA 251 	ATATCGAGCTAAGAAAATAATATTAGCTTACGCTTTGGCTGTAAGTGGTATTCATAATGT 191	72 AGTACGTATACTAAACTTTAAGCAAACAGATGAGGGTCATACAGCATCATATGGAATTGA 131 	12 ATTAACACCTGATGATAGAAGTGGATATCCACCCGTTGAAAAGCAAATAGCAGGAGATAT 71	vative 0; Mismatches 162; Indels 54; Gaps 2;

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Sequence 5, Application US/09463048A

Patent No. 6630619

GENERAL INFORMATION:

APPLICANT: Commonwealth Scientific and Industrial Research Organisation
APPLICANT: Commonwealth Scientific and Industrial Research Organisation
APPLICANT: EAST, Peter David

TITLE OF INVENTION: Toxin Genes from the Bacteria Xenorhabdus nematophilus
TITLE OF INVENTION: Uninsescens

FILE REFERENCE: 050179-0076

CURRENT APPLICATION NUMBER: US/09/463,048A

CURRENT ETLING DATE: 2002-12-13

PRIOR APPLICATION NUMBER: PCT/AU98/00562

PRIOR APPLICATION NUMBER: PO8088

PRIOR FILING DATE: 1998-07-17

PRIOR APPLICATION NUMBER: 20

NUMBER: OF SEQ ID NOS: 20

COETMADE: TOXING DATE: 1997-07-17
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; SEQ ID NO 5
; LENGTH: 1205
; TYPE: DNA
; ORGANISM: Xenorhabdus nematophilus
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Best Local Similarity
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                  TTATATTGAGAGAAAAAAACTTTCTAAACCTTTCTTTCGACCACCGCAGAGATTACCTGA
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Pred. No. 2.7e-157;
0; Mismatches 162;
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; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA (gen
US-08-569-168-6
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  Query Match 62.3%;
Best Local Similarity 79.1%;
Matches 833; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Smigielski, Adam J.
APPLICANT: Akhurst, Raymond J.
TITLE OF INVENTION: TOXIN GENE
NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IEM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIN Release #1.0, VC

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/569,168
                                                                                                                                                                                                                                                                   CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Price, Robert L.
REGISTRATION NUMBER: 22,685
REFERENCE/DOCKET NUMBER: 1451
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                           TELEPHONE: 703-684-1111
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U
ZIP: 22314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Alexandria
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                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                       LENGTH:
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99 Canal Center Plaza, Suite 300
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Score 628.2; DB 2;
Pred. No. 2.5e-148;
0; Mismatches 163;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Version
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                        CATAGTGTTGGAACTCATTATAAAAATATATGA 1008
                                                                                          CTAAGCCAAAAAGACAGGGTAACCTTTACTCAAAATGATACTGACAATACAATGACGGTT
                                                                                                                                                           GGTACAGCCAGTGGCAGTAACCTGGAAAAATTGCCGAATA---ATTTATATAGTGTGAGG
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CAAAGTGTTGGAACTCATTACCAAAATATATAA
                                                                     CTAAGCCAAGAACACAGGGTAACATTCTCCATAAATAATACTGACCAAATAATGGAGATC
                                                                                                                                          AGTTCATTTATAAAAGAGGTACAAAAGGGTAATGATCCACGTGCAGCAGCAGCAGCAAGTATT
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FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Price, Robert L.
REGISTRATION NUMBER: 22,685
REFERENCE/DOCKET NUMBER: 1451
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-684-1111
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 604; Conserv
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APPLICANT: Smigjelski, Adam J.
APPLICANT: Akhurst, Raymond J.
APPLICANT: Akhurst, Raymond J.
TITLE OF INVENTION: TOXIN GENE FROM XENORHABDUS NEMATOPHILUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 22314
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: 1i MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lowe, Price, Leblanc & Becker STREET: 99 Canal Center Plaza, Suite 300
CITY: Alexandria
STATE: VA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
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TTGGAATGTTCCTGAGCTTGAAAATAAACCATT-GAGTGCTTATTCAGATGACGATAAAT
                                          AATACGTGTACTAGAATTCAAGCAAACAAATGAAAGTCATACAGGATTGTATGGAATTCC
                                                                                                            ATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAAATTTCTAAAGA
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                                                                                                                                                  Query Match
Best Local Similarity
Matches 604; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Smigielski, Adam J.
APPLICANT: Akhurst, Raymond J.
TITLE OF INVENTION: TOXIN GENE FROM XENORHABDUS NEMATOPHILUS
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-684-1111
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                             TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 837 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Price, Robert L.
REGISTRATION NUMBER: 22,685
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: >> _ CITY: Alexandria
                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
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AATACGTGTACTAGAATTCAAGCAAACAAATGAAAAGTCATACAGGATTGTATGGAATTCC 176
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99 Canal Center Plaza, Suite 300
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                                                                                                                                                Score 521.4; DB 2;
Pred. No. 1.6e-121;
0; Mismatches 101;
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                                 8
                                                                                                                                                ; NAME/KEY: promoter
; LOCATION: (1)..(1141)
; OTHER INFORMATION: consensus sequence
US-09-806-708B-22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22, Application US/09806708B Patent No. 6784342 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.0 SEQ ID NO 22 LENGTH: 1141
                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: The University of British Columbia TITLE OF INVENTION: Regulation of Embryonic TrILE REFERENCE: 4810-58741
CURRENT APPLICATION NUMBER: US/09/806,708B
CURRENT FILING DATE: 2001-04-03
                                                                                                                                                                                                                                                                                                                                                    CURRENT FILING DATE: 2001-04-03
PRIOR APPLICATION NUMBER: US 60/147,133
PRIOR FILING DATE: 1999-08-04
                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                         ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                             TYPE: DNA
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                                   185 ATAATGTATCTAAACTTCCTGATGACTATTATAAGAATAAAGAGACTGCTGAGAGAATTT 244
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                                                                          81;
                                                                                         Similarity
ACRNTRTWWABWKHSWCNNNNNNNNNNNNNTWCHYTTANABBCYRANNNNAAARMARTC 775
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                                                                                         6.2%;
                                                                      Score 62.6; DB 4;
Pred. No. 4.4e-06;
0; Mismatches 360;
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                                                                                                                                                                                                                                                                                                                                                   Sequence 14, Application US/08232463 Patent No. 5670367
                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
          COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                         APPLICANT: SCHEIFLINGER, F.
APPLICANT: FALKNER, F.G.
TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
NUMBER OF SEQUENCES: 52
                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                  APPLICANT: DORNER, F.
                                                                                                                                                                      CITY: Alexandria
APPLICATION NUMBER:
                                                                                                                     COUNTRY: USA
ZIP: 22313-0299
                                                                                                                                                        STATE:
                                                                                                                                                                                          STREET:
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US/08/232,463
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NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 304*
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: line IMMEDIATE SOURCE:
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PRIOR APPLICATION NUMBER: 1
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TYPE: nucleic acid
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 6.0%;
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1124
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ACAATGGATTAGAGGCTAAATCATTAGAGAATTATATTGAGAGAAAAAAACTTTCTAAAC
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                         TGGAAAAATTGCCGAATAATTTATATAGTGTGAGGCTAAGCCAAAAAAGACAGGGTAA 937
                                                                                                                ANAAGGGTAACGCTCCACAGACAGCAGCGAAAAGTATTGGTACAGCCAGTGGCAGTAACC
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5.0%; Pred. No. 4.3e-05;
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US-09-806-708B-23
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Patent No. 6784342
GENERAL INFORMATION:
APPLICANT: The University of British Columbia
TITLE OF INVENTION: Regulation of Embryonic Transcription
FILE REFERENCE: 4810-58741
CURRENT APPLICATION NUMBER: US/09/806,708B
CURRENT FILING DATE: 2001-04-03
PRIOR APPLICATION NUMBER: US 60/147,133
PRIOR FILING DATE: 1999-08-04
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn version 3.0
SEQ ID NO 23
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US-09-806-708B-22
; Sequence 22, Application US/09806708B
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LOCATION: (1)..(1055)
OTHER INFORMATION: consensus sequence of A.t. and L.a. FAE1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 1055
TYPE: DNA
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                                                                                  WTCKAWMKAWATGAATTTNA
                                                                                                               TCATTTATAAAAGAGGTACA 821
                                                                                                                                                                                                             YTTRTANNGACTTTTTNNTTGGMRTNTAAARGWANNNNNNNNNNNNNNNGACWAWRTTTAT
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                                                                                                                                             ANCGTNINININININAYATTTNTATTTTWKTKKANININININAAAYYGAAAWKUNTTMC
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Pred. No. 0.0035;
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; SEQ ID NO 22
; LENGTH: 1141
; TYPE: DNA
; ORGANISM: Artificial sequence
; FEATURE:
; NAME/KEY: promoter
; LOCATION: (1)..(1141)
; OTHER INFORMATION: consensus sequence of US-09-806-708B-22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Patent No. 6784342; GENERAL INFORMATION:
; GENERAL INFORMATION:
; APPLICANT: The University of British Columbia; TITLE OF INVENTION: Regulation of Embryonic Transfille OF INVENTION: Regulation of Embryonic Transfille OF INVENTION: REGULATION NUMBER: US/09/806,708B; CURRENT FILING DATE: 2001-04-03; PRIOR APPLICATION NUMBER: US 60/147,133; PRIOR FILING DATE: 1999-08-04; NUMBER OF SEQ ID NOS: 23; SOFTWARE: PatentIn version 3.0
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5.0%; Score 50.6; D
Local Similarity 11.5%; Pred. No. 0.00
nes 101; Conservative 313; Mismatches
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715 NNAGTWKNNNNNAKNASAAKNYÄÄÄAVKÄÄKKHWRWANKWAMRGWHÄDÄÄÄBTTDKRNN
                                                                                                                                                                                                                                                                                                                                                                                          524 TATTTACCTTATCTGCTGTAACTTCACTGGGAAGGAGGATTTTTTCAAAAAACTTTTACA
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                                         TTTCTAATATGGAAGGGGCTGCAAAACAAAAGTATAGTTCATTTATAAAAGAGGTACAAA 823
                                                                                                                           AAGCGCCTAAATGGAGAGTGAGTTTTAAAGAACTTAAAAATAACAAATCTAGGAATGGAT
                                                                                                                                                                        BWSHTWBHBRAGAAHYWMBMMYBAKCHCMKAWYKAKKYAGAGGSNNNNNNNNNNNNNNNNNNN
                                                                                                                                                                                                                 TCTTTCGACCACCGCAGAGATTACCTGATGGCAGAATAGGTTATTTGGCTGGACCAACAG
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                                                                                    ATCARDDYYAASRWYAMANAKWYYYKBAANNAYYTHANNWWGCWNNATDTRRTMWKNNNN
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US-09-601-198-153
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Best Local Similarity 47.3%;
Matches 249; Conservative
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: NUCLEIC ACID PROBES AND METHOD FOR DETECTING UREAPLASMA FILE OF INVENTION: UREALYTICUM FILE REFERENCE: UAB-13452/22 CURRENT FAILOR NUMBER: US/09/601,198 CURRENT FILING DATE: 2000-12-08 PRIOR APPLICATION NUMBER: 60/073,189 PRIOR APPLICATION NUMBER: 60/073,189 PRIOR FILING DATE: 1998-01-30
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Glass, John I.
Heiner, Cheryl R.
Lefkowitz, Elliot
ATCTGCTGTAACTTCACTGGGAAGGAGGATTTTTTCAAAAAACTTTTACAATGGATTAGA
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                                                                                                    --ATTTATCAAGAATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATC
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                                                              ATGACGATAAATT-----ATTAGCACTATATTTTTTCTCTGTACAGGAAATTCCACTGGA 473
                                                                                                                                                                                                                           AATTAATAGAATTTAGTAATTATTTTAAAAACCAAAAAACCACCAAGAGATTCTAATGAAT 1336
                                                                                                                                                                                                                                                               ATCCTCAAAACATTTGGAATGTTCCTGAGCTTGAAAATAAACCATTGAGTGCTTATTCAG 418
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                                                                                                                                            ATAAGCAAAAATTAAATGATTATGAAAAAATATTTTCACAAAATTGGTATTTTATTTTAA
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Fllson Y.
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Pred. No. 0.016;
0; Mismatches 267;
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; FEATURE:
; NAME/KEY: promoter
; LOCATION: (1)..(1055)
; OTHER INFORMATION: consensus sequence of A.t.
US-09-806-708B-23
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; ORGANISM: Staphylococcus epidermidis
US-09-134-001C-1161
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US-09-134-001C-1161
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APPLICANT: The University of British Columbia
TITLE OF INVENTION: Regulation of Embryonic T-
FILE REFERENCE: 4810-58741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 5674
SEQ ID NO 1161
LENGTH: 729
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                                                                                                                                                                                                                                                                                                                                                                Sequence 23, Application US/09806708B Patent No. 6784342
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                                                                                                                                                      SEQ ID NO 23
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Best Local Similarity
                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/134,001C CURRENT FILING DATE: 1998-08-13 PRIOR APPLICATION NUMBER: US 60/064,964
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                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/806,708B CURRENT FILING DATE: 2001-04-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 1997-11-08
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: GTC-007
                                                                                                                                                                          SOFTWARE:
                                                                                             LENGTH: 1055
TYPE: DNA
ORGANISM: Artificial sequence
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                                                                                                                                                                          PatentIn version 3.0
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                                                                                                                                                                                                                                                                                                           Transcription in Plants
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                       and L.a.
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                   FAE1 promoters
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RESULT 14
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                                                                                                                                                                               Sequence 531, Applicat Patent No. 6593114
GENERAL INFORMATION:
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Best Local
NUMBER OF SEQUENCES: 5256
CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences,
                                       Patrick S. Dillon
Craig A. Rosen
Steven C. Barash
Michael R. Fannon
TITLE OF INVENTION: Staphylococcus
NUMBER OF SEQUENCES: 5256
                                                                                                                                                                 APPLICANT: Charles Kunsch
                                                                                                                                                                                                                                                                                                                                                                              104
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                                                                                                                                                                                                                                                                                                                                          AGAACTTAAAAATAACAAATCTAGGAATGGATTTTCTAAT 771
                                                                                                                                                                                                                                                                                                                                                                            MYTAGKAAWTNNKTASGKWMYAMMKTTWYWAYWTWCAACWMATAKRRTAAKMCWYYTRGY
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                                                                                                                                                                                                                                                                                                                                                                                                           TWSTCRWKTARGWWYYAMRYYTWAK-ASGCNNNAKWRCAWWYACATNNNAMAWWYACAYW
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                                                                                                                                                                                                                    Application US/08956171E
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                                                                                                                                                 Choi
                                                             aureus
                                                             Polynucleotides
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                                                           Sequences
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Sequence 531, Application U.
Patent No. 6737248
GENERAL INFORMATION:
APPLICANT: Charles Kunson TITLE OF INVENTION: Star
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Best Local Similarity
Matches 138; Conserv
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TELEFAX: (301) 3(
INFORMATION FOR SEQ ID NO:
                                                                CORRESPONDENCE ADDRESS:
                                                                                  NUMBER OF SEQUENCES:
CITY:
                  STREET:
                                             ADDRESSEE:
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FILING DATE: January 5, 1996
APPLICATION NUMBER: 08/781,986
FILING DATE: January 3, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Mark J. Hyman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS Version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 46,789
REFERENCE/DOCKET NUMBER: PB248P1
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                     306 TAAAGATATGGCAAATGGTTTTTATAAGAATGAACTGGATTTTGAAGGT
                                                                                                                                                                                                                                                                                                         553 AAAACAAGGAACATACGCTTTTATTAAGAAAGAAGATTCTATTAAAGAT
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: 9410 Key West Avenue Rockville
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TYPE: nucleic acid
STRANDEDNESS: double
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CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: Maryland
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                                         Human
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47.8%;
                                             Genome Sciences,
                                                                                                                                  Kunsch
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                                                                                    Staphylococcus aureus 5255
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Pred. No. 0.027;
0; Mismatches 151;
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COUNTRY: USA
ZIP: 20850
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/781,986A
FILING DATE:
CLASSIFICATION NUMBER: BELICATION:
APPLICATION NUMBER: PILING DATE:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Benson, Bob
REGISTRATION NUMBER: PB248PP
TELEPHONE: 30,446
REFERENCE/DOCKET NUMBER: PB248PP
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
TELEPAX: (301) 309-8504
TELEPAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: 531:
SEQUENCE CHARACTERISTICS:
LENGTH: 942 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
US-08-781-986A-531

4.7%; Score 47.4; DB 4;
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Search completed: March
Job time : 221 secs
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                                                                                                553
                                                                                                                                      306 TAAAGATATGGCAAATGGTTTTTATAAGAATGAACTGGATTTTGAAGGT 354
                                                                                                                                                                                              493 AACGCCATCAAATGCTGATTTTTGTATTTTAGTTAATGGTGATTCAATGGAACCTATGTT 552
                                                                                                                                                                                                                                           246 TCAAGAATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAAATTTC 305
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                         1, 2005, 06:54:52
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Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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ALIGNMENTS

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Sequence 2, Application US/10617962

Sequence 2, Application US/10617962

GENERAL INFORMATION:

APPLICANT: Commonwealth Scientific and Industrial Research Organisation
APPLICANT: EAST, Peter David

TITLE OF INVENTION: Toxin Genes from the Bacteria Xenorhabdus nematophilus and Photor

TITLE OF INVENTION: luminescens

FILE REFERENCE: 050179-0076

CURRENT APPLICATION NUMBER: US/10/617,962

CURRENT APPLICATION NUMBER: US/09/463,048A

PRIOR APPLICATION NUMBER: US/09/463,048A

PRIOR APPLICATION NUMBER: PCT/AU98/00562

PRIOR PILING DATE: 1998-07-17

NUMBER OF SEQ ID NOS: 20

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; LENGTH: 1008
; TYPE: DNA
; ORGANISM: Photorhabdus luminescens
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                                                                     CCTCAAAACATTTGGAATGTTCCTGAGCTTGAAAATAAACCATTGAGTGCTTATTCAGAT
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                       CAACAATCAAATGCCGCAAGATTTTTTAAATTAATTGATTTCTTATTTACCTTATCTGCT
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CURRENT APPLICATION NUMBER: US/10/617,962
CURRENT FILING DATE: 2003-07-14
PRIOR APPLICATION NUMBER: US/09/463,048A
PRIOR FILING DATE: 2002-12-13
PRIOR APPLICATION NUMBER: PCT/AU98/00562
PRIOR APPLICATION NUMBER: PCT/AU98/00562
PRIOR FILING DATE: 1998-07-17
PRIOR APPLICATION NUMBER: PO 8088
PRIOR FILING DATE: 1997-07-17
PRIOR APPLICATION NUMBER: PCT/AU98/00562
PRIOR FILING DATE: 1998-07-17
PRIOR APPLICATION NUMBER: PCT/AU98/00562
PRIOR FILING DATE: 1998-07-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1, Application US/10617962 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Xenorhabdus nematophilus -10-617-962-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Commonwealth Scientific and Industrial Research Organisation
APPLICANT: EAST, Peter David
TITLE OF INVENTION: Toxin Genes from the Bacteria Xenorhabdus nematophilus and
TITLE OF INVENTION: luminescens
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0; Mismatches 162;
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TITLE OF INVENTION: luminescens
FILE REFERENCE: 050179-0076
CURRENT APPLICATION NUMBER: US/10/617,962
CURRENT FILING DATE: 2003-07-14
PRIOR APPLICATION NUMBER: US/09/463,048A
PRIOR FILING DATE: 2002-12-13
PRIOR APPLICATION NUMBER: PCT/AU98/00562
PRIOR FILING DATE: 1998-07-17
PRIOR APPLICATION NUMBER: PCT/BUSS/00562
PRIOR APPLICATION NUMBER: PO 8088
                                                                                                                                                                                                                                                                   Sequence 5, Application US/10617962 GENERAL INFORMATION:
                                                                                                                                                                                                                             APPLICANT: Commonwealth Scientific and Industrial Research Organisation APPLICANT: EAST, Peter David
PRIOR FILING DATE: 1997-07-17
NUMBER OF SEQ ID NOS: 20
SOFTWARE: PatentIn version 3.
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TACAAAAAGCGGCAGTAACTTCGAAAAACTGCAAGGTAGAGATTTATATAGATATAAGACT
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Pred. No. 1.5e-135;
0; Mismatches 162;
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Sequence 14, Application US/08466194
GENERAL INFORMATION:
APPLICANT: DORNER, F.
APPLICANT: SCHEIFLINGER, F.
APPLICANT: FALKNER, F. G.
TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & Lardner
STREET: 3000 K Street, NW, Suite 500
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US-08-466-194-14/c
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US-08-466-194-14
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                                                                                                                                 Query Match
Best Local S
Matches 25
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SEQUENCE CHARACTERISTICS:
LENGTH: 7218 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 22-APR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/935
FILING DATE: 26-AUG-1992
PRIOR APPLICATION NUMBER: EP 91 114
APPLICATION NUMBER: EP 91 114
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 3047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve.
CURRENT APPLICATION NUMBER: US/08/466,194
FILING DATE: 06-JUN-1995
PRIOR APPLICATION NUMBER: US/08/232,463
                                                                                                                                                                                                                                  IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: I
                                                                                                                               y Match
5.9%; Score 59.8; DB 10;
Local Similarity 6.0%; Pred. No. 0.027;
hes 25; Conservative 225; Mismatches 167;
                                                                                                                                                                                                                                                    STRANDEDNESS: single TOPOLOGY: linear
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CITY: Washington
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                                                                                                TCTTATTTACCTTATCTGCTGAACTTCACTGGGAAGGAGGATTTTTTCAAAAAACTTTT
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(202)672-5399
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RESULT 5
PCT-US99-26796-234
Sequence 234, Application PC/TUS9926796
GENERAL INFORMATION:
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Best Local S
Matches 272
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SEQ ID NO 234
LENGTH: 3927
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CURRENT FILING DATE: 1999-11-05
NUMBER OF SEQ'ID NOS: 420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 43.9
mes 272; Conservative
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                                                                                                                                         TAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAAATTTCTAAAGATATGGCAAA
                                                                                                                                                                                                              GAMAATAATTAGCTTACGCTTTGGCTGTAAGTGGTATTCATAATGTATCTAAACTTCC
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TGTTGAAAATTTATTGAAAGAACACAAAATAGCTGTTAATGAAAGTGAAGAAATAGCACA
                                                                 TGGTGATAATATAGATGATGATAATATATATGATGGTGATAACATAAATAGTGGTGATAA 2870
                                                                                                   TGGTTTTTATAAGAATGAACTGGATTTTGAAGGTCAATATCCTCAAAACATTTGGAATGT 380
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 52.6; DB 1;
Pred. No. 0.87;
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; TYPE: DNA
; CTGANISM: Homo sapiens
; FEATURE:
; FANTURE:
; LOCATION: (1)...(703368)
; OTHER INFORMATION: n = A,T,C or
US-10-990-328-94720
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; Sequence 94720, Application US/10990328
; GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 5
SOFTWARE: FastSEQ for W
SEQ ID NO 94720
LENGTH: 709368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: CARGILL, M
TITLE OF INVENTION: F
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 169;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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CURRENT APPLICATION NUMBER: US/10/990,328
CURRENT FILING DATE: 2004-11-17
674225 ATTAAGAGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Match 5.2%;
Local Similarity 45.7%;
                                                                                                                                                            674345 GATGGAATGCGAGAATAAAGCTGAATTTATCTATKTTCAYTTATGATTAAAAAAAAGATAA
                                                                                                                                                                                                                                                                                                                                                                                                       674525 TAACAATAAATATTCCATTTTAATAGTATGTTAAAAAAATCACTCATAAGCCTTTCCTTTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          674585 TATCTCATCACACAAATCAAAAATTAGGAAATGTAGTAGCAYATTTCACATGATCTTTTG
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                                                                                                                                                                                                                                                                               629 AACTITCTAAACCTTTCTTTCGACCACCGCAGAGATTACCTGATGGCAGAATAGGTTATT
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                                        TAAAAGAGGT
                                                                                                                AATCTAGGAATGGATTTTCTAATATGGAAGGGGCTGCAAAACAAAAGTATAGTTCATTTA
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                                                                                ATTCTAAGTAGGTATTTTAAAAGATAGTTTGACATATACACTCCCAGTGTTGGACTTGAA
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POLYMORPHISMS IN NUCLEIC ACID MOLECULES
ENCODING HUMAN ENZYME PROTEINS, METHODS OF DETECTION
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Pred. No. 4.9;
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RESULT 8
US-10-235-192A-27
; Sequence 27, Application US/10235192A
; Sequence 27, Application US/10235192A
; GENERAL INFORMATION:
; APPLICANT: McCarthy, Jeanette
; TITLE OF INVENTION: Methods and Compositions for Identifying
; TITLE OF INVENTION: And Disorders for Abnormal Lipid Levels a
; TITLE OF INVENTION: and Disorders Associated Therewith
; FILE REFERENCE: MMI-011
; CURRENT APPLICATION NUMBER: US/10/235,192A
; CURRENT FILING DATE: 2002-09-04
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US-10-473-126-386/c
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LENGTH: 8056
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/473,126
CURRENT FILING DATE: 2003-09-26
CURRENT FILING DATE: 1258
NUMBER OF SEQ ID NOS: 1258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Epigenomics AG
TITLE OF INVENTION: Methods and nucleic acids for the
TITLE OF INVENTION: proliferative disorders
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SEQ ID NOS:
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Pred. No. 2.4;
0; Mismatches 270;
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NAME/KEY: misc feature
LOCATION: 145329-145428, 187896-187995,
OTHER IMPORMATION: N = any nucleotide
US-10-235-192A-27
                                                                                                                                                              PCT-US99-26796-311
FCT-US99-26796-311
FSQUENCE 311, Application PC/TUS9926796
GENERAL INFORMATION:
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Matches
                        APPLICANT: United States STATES APPLICANT: United States STATES APPLICANT: Secretary of the Navy TITLE OF INVENTION: Chromosome 2 Sequence of Human Malaria Parasite TITLE OF INVENTION: Plasmodium Falciparum and Proteins of Said TITLE OF INVENTION: Chromosome Useful in Anti-malarial Vaccines and TITLE OF INVENTION: Diagnostic Reagents
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SEQ ID NO 27
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FILE REFERENCE:

CURRENT APPLICATION NUMBER: PCT/US99/26796

TITING DATE: 1999-11-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo
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Local Similarity 42.6%;
hes 263; Conservative
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APPLICANT: GOFF, STEPHEN
APPLICANT: BONNA, CAROLINE
APPLICANT: BONNA, CAROLINE
APPLICANT: COLBERT, MICHELLE
APPLICANT: WANG, RONG-LIN
TITLE OF INVENTION: CEREAL TRINUCLEOTIDE SIMPLE SEQUENCE
TITLE OF INVENTION: REPEAT MARKERS AND THEIR USES
FILE REFERENCE: NADII 0.58C1
CURRENT APPLICATION NUMBER: US/10/266,090
CURRENT APPLICATION NUMBER: US 10/260,703
PRIOR APPLICATION NUMBER: US 00/326,117
PRIOR APPLICATION NUMBER: US 60/326,117
                                                                                                                                                                                                           RESULT 10
US-10-266-090-23096
; Sequence 23096, Application
; GENERAL INFORMATION:
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; SOFTWARE: Patentin Ver
; SEQ ID NO 311
; LENGTH: 3738
; TYPE: DNA
; ORGANIAM: Plasmodium f
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Best Local Similarity
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Pred. No. 2.6;
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; PRIOR FILING DATE: 2001-09-26; NUMBER OF SEQ ID NOS: 51812; SOFTWARE: FastSEQ for Windows Ve; SEQ ID NO 23096
; LENGTH: 1145
; TYPE: DNA; ORGANISM: ZEA MAYS; FEATURE: MAME/KEY: misc feature; LOCATION: (1)... (1145); OTHER INFORMATION: n = A,T,C or US-10-266-090-23096
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                                                 TCCACAGACAGCAGCGAAAAGTATTGGTACAGCCAGTGGCAGTAACCTGGAAAAATTGCC
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                                                                                               ACCGCAGAGATTACCTGATGGCAGAATAGGTTATTTGGCTGGACCAACAGAAGCGCCTAA 713
                                                                                                                                                                                                                                                                                                                        AATATCCTCAAAACATTTGGAATGTTCC--TGAGCTTGAAAATAAACCATTGAGTGCTTA 413
GAATAATTTATATAGTGTGAGGCTAAGCCAAAAAGACAGGGTAACCTTTACTCAAAATGA
                                                                                                                                                                      GGAAGGGGCTGCAAAACAAAAGTATAGTTCATTTATAAAAGAGGTACAAAAGGGTAACGC
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RESULT 11 US-10-240-453-95 ; Sequence 95, A)

Application US/10240453

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                                        ; OTHER INFORMATION: n = US-09-948-124-61
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US-09-948-124-61
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                                                                                                                                                                                                                                          Sequence 61, Application US/09948124
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig
TITLE OF INVENTION: COLLECTION OF SINGLE NUCLEOTIDE POLYMORPHISM (SNPs) LOCATED ON
TITLE OF INVENTION: Y, METHODS OF DETECTION, AND USES THEREOF
FILE REFERENCE: CL001301
CURRENT APPLICATION NUMBER: US/09/948,124
CURRENT FILING DATE: 2002-12-23
                                                                                                                                                                               NUMBER OF SEQ ID NOS:
SEQ ID NO 61
LENGTH: 318007
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Best Local (
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LENGTH: 11394
  Query Match
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APPLICANT: OLEK, Alexander
APPLICANT: PIEDENBROCK, Christian
APPLICANT: BERLIN, Kurt
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CURRENT APPLICATION NUMBER: US/10/240,453
CURRENT FILING DATE: 2002-10-02
PRIOR APPLICATION NUMBER: PCT/EP01/03973
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: DE 10043826.1 PRIOR FILING DATE: 2000-09-01
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PRIOR APPLICATION NUMBER: DE 10019058.8
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                                                                              NAME/KEY: misc_feature
LOCATION: (1)...(31800)
                                                                                                                                         ORGANISM: HUMAN
                                                                                                                                                                TYPE: DNA
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                                                                                                                        FEATURE:
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Pred. No. 4.8;
  Score 49.8;
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US-09-947-911-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 2
LENGTH: 600322
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                                                                                                                                                                                                                                                                                Matches 159;
                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: VENTER, J. Craig
TITLE OF INVENTION: COLLECTION OF SINGLE NUCLEOTIDE POLYMORPHISM (SNP8) LOCATED ON
TITLE OF INVENTION: X, METHODS OF DETECTION, AND USES THEREOF
FILE REFERENCE: CL001300
CURRENT APPLICATION NUMBER: US/09/947,911
CURRENT FILING DATE: 2001-09-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF
                                                                                                                                                                                                                                                                                                                                                    FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(600322)
OTHER INFORMATION: n = A,T,C or
                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: HUMAN
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Pred. No. 1
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Sequence 238, Applicat GENERAL INFORMATION:
APPLICANT: United St APPLICANT: Secretary TITLE OF INVENTION:
TITLE OF INVENTION:
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PCT-US99-26796-238
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LENGTH: 5940
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Best Local 9
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SOFTWARE: PatentIn V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: PCT/US99/26796
CURRENT FILING DATE: 1999-11-05
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                                GAAGGGGCTGCAAAACAAAGTATAGTTCATTTATAAAAGAGGTACAAAAGGGTAA 830
                                                                                                                                                                                                     TGAAAAAGAAAAAAAAAAAAAAGGAAGAAACATTTCATAATATAGAAAAAGAGTA
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                                                                                              TGGAGAGTGAGTTTTAAAGAACTTAAAAAATAACAAATCTAGGAATGGATTTTCTAATATG
                                                                                                                                   TTTAGAAAAAATAAAGAAAGAAACGATTTCTATAGAAATTATAGATATTAAAAAACA
                                                                                                                                                                    CCGCAGAGATTACCTGATGGCAGAATAGGTTATTTGGCTGGACCAACAGAAGCGCCTAAA
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Secretary of the Navy
VENTION: Chromosome 2 Sequence of Human Malaria Parasite
VENTION: Plasmodium Falciparum and Proteins of Said
VENTION: Chromosome Useful in Anti-malarial Vaccines and
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Pred. No. 5.5;
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US-09-865-419A-12520
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APPLICANT: Wu, Kunsheng
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(51935)B
CURRENT APPLICATION NUMBER: U5/09/865,419A
CURRENT FILING DATE: 2001-05-29
PRIOR APPLICATION NUMBER: US 60/208,063
PRIOR APPLICATION NUMBER: US 60/208,063
PRIOR FILING DATE: 2000-05-31
NUMBER OF SEQ ID NOS: 54020
SEQ ID NO 12520
LENGTH: 572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: unsure
LOCATION: (1)..(572)
COTHER INFORMATION: unsure at all n locations
OTHER INFORMATION: Clone ID: LIB3236-144-P1-N1-H3
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ORGANISM: Zea
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554
                                                                                                  870 TGGCAGTAACCTGGAAAAATTGCCGAATAATTTATATAGTGTGAGGCTAAGCCAAAAAGA
                                                                                                                                      314 АЗАЗДАЗАЗАЗДАЗАЗАЗДАЗТТТТТЯЗАТТТТТАЗАЗДАЗАЗАЗАЗАЗАЛИТАЗАЗАЗАЗАЗА
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TAAAAAAANTTATT
                                CAGGGTAACCTTTACT 945
                                                                   AAAAGAGGTACAAAAGGGTAACGCTCCACAGACAGCGGCGAAAAGTATTGGTACAGCCAG
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569
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Pred. No. 3.2;
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Search completed: March 1, 2005, 08:11:55 Job time: 4616 secs

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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen

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Result
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Maximum
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Maximum Match 100%
Listing first 45 summaries
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         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 2000000000
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1: /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*

2: /cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*

3: /cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*

4: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*

5: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*

6: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*

7: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*

8: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*

9: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*
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6 US-10-932-182A-76518

7 US-10-517-441-618

8 US-10-517-441-68

8 US-11-044-879-3

7 US-10-450-763-12549

10S-10-932-112A-6042

7 US-10-517-441-293

17 US-10-517-441-293

17 US-10-517-441-567

17 US-10-517-441-567

17 US-10-517-441-567

17 US-10-517-441-581

17 US-10-517-441-581

17 US-10-517-441-581

17 US-10-517-441-568
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US-10-991-672A-4

5 US-10-954-778-160

5 US-10-954-778-160

7 US-10-517-441-307

7 US-10-517-441-715

7 US-10-932-182A-166164
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Sequence 166160,
Sequence 4, Appli
Sequence 160, App
Sequence 307, App
Sequence 307, App
Sequence 715, App
Sequence 715, App
Sequence 1, Appli
Sequence 43, Appli
Sequence 44, App
Sequence 688, Appli
Sequence 688, Appli
Sequence 688, Appli
Sequence 684, Appli
Sequence 689, Appli
Sequence 581, Appli
Sequence 584, Appli
Sequence 584, Appli
Sequence 5042, Appli
Sequence 567, Appli
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Sequence 317, App	Sequence 75787, A Sequence 1492, Ap	Sequence 293, App Sequence 567, App	Sequence 1, Appli Sequence 166179,	Sequence 442, App		_	Sequence 37, Appl	Sequence 738, App			Sequence 41, Appl		b	•	Sequence 322, App	Sequence 737, App

ALIGNMENTS

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US-10-932-182A-166160
; Sequence 166160, Application US/10932182A
; GENERAL INFORMATION:
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; ORGANISM: Saccharomyces pastorianus
US-10-932-182A-166160
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SEQ ID NO 166160
LENGTH: 1134
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Best Local Similarity
Matches 144; Conserv
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APPLICANT: NAKAMURA, NORIHISA
APPLICANT: WODAMA, YUKIKO
APPLICANT: FUJIMURA, TOKHIHIKO
APPLICANT: FUJIMURA, TOKHIHIKO
TITLE OF INVENTION: METHODS FOR ANALYZING GENES OF INDUSTRIAL YEASTS
FILE REFERENCE: 030685-043
CURRENT APPLICATION NUMBER: US/10/932,182A
CURRENT FILING DATE: 2004-09-02
NUMBER OF SEQ ID NOS: 197023
                                                                                                                                                                                                                                                                                                                               610
850
                                                                                                                    328 TATAAGAATGAACTGGATTTTGAAGGTCAATATCCTCAAAACATTTGGAATGTTCCTGAG
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TTATATAATAATAATATTGATAATATTCCTATAGATATTTTAATATTTAAATATTT
                                CTTGAAAATAAACCATTGAGTGCTTATTCAGATGACGATAAATTATTAGCACTATATTT
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RESULT 2

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RESULT 3
US-10-954-778-160/c
Sequence 160, Application US/10954778
GENERAL INFORMATION:
APPLICANT: ZWIEBEL, LAURENCE J.
TITLE OF INVENTION: METHOD OF IDENTIFYING
TITLE OF INVENTION: ODORANT RECEPTORS OF
FILE REFERENCE: N1125
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CURRENT APPLICATION NUMBER: US/10/954,778
CURRENT FILING DATE: 2004-09-30
PRIOR APPLICATION NUMBER: 10/056,405
PRIOR FILING DATE: 2002-01-24
PRIOR APPLICATION NUMBER: 60/264,649
PRIOR APPLICATION NUMBER: 60/264,649
PRIOR FILING DATE: 2001-01-26
NUMBER OF SEQ ID NOS: 248
SOFTWARE: Patentin Ver. 3.3
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LENGTH: 1941
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/691,672A
CURRENT FILING DATE: 2003-10-24
NUMBER OF SEQ ID NOS: 13
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Pred. No. 0.12;
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US-10-954-778-160
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                                                                                                                                                                                                                                                        SEQ ID NO 160
LENGTH: 3089
TYPE: DNA
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LENGTH: 3089
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Best Local Similarity 44.3%;
Matches 185; Conservative
                                                                                                                                             Matches 191;
                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                             APPLICANT: ZWIEBEL, LAURENCE J.
TITLE OF INVENTION: METHOD OF IDENTIFYING CHEMICAL AGENTS
TITLE OF INVENTION: ODORANT RECEPTORS OF SENSORY NEURONS
FILE REFERENCE: N1125
FILE REFERENCE: N1125
CURRENT APPLICATION NUMBER: US/10/954,778
CURRENT FILING DATE: 2004-09-30
PRIOR APPLICATION NUMBER: 10/056,405
PRIOR FILING DATE: 2002-01-24
PRIOR FILING DATE: 2002-01-24
PRIOR APPLICATION NUMBER: 60/264,649
PRIOR APPLICATION NUMBER: 60/264,649
PRIOR APPLICATION NUMBER: 2001-01-26
NUMBER: 05 SEQ ID NOS: 248
NUMBER: 05 SEQ ID NOS: 248
                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Ver. 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Anopheles gambiae
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288
                                  139 GCTAAGAAAATAATTAGCTTACGCTTTGGCTGTAAGTGGTATTCATAATGTATCTAAA 198
                                                                                                          ATACTAAACTTTAAGCAAACAGATGAGGGTCATACAGCATCATATGGAATTGAATATCGA 138
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TGAAGGTCAATATCCTCAAAACATTTGGAATGTTCCTGAGCTTGAAAATAAACCATTGAG 407
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                                                                                                                                               Conservative
                                                                                                                                                                4.3%;
43.7%;
                                                                                                                                                                                                                                         gambiae
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                                                                                                                                             0;
                                                                                                                                             Score 43.4; DB
Pred. No. 0.52;
0; Mismatches :
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                                                                                                                                                                                 DB 6;
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                                                                                                                                               246;
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US-10-517-441-307
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PRIOR APPLICATION NUMBER: PCT/EP2003/010881
PRIOR FILING DATE: 2003-10-01
PRIOR APPLICATION NUMBER: DE 10317955.0
PRIOR FILING DATE: 2003-04-17
PRIOR FILING DATE: 2003-04-17
PRIOR APPLICATION NUMBER: DE 1030096.8
PRIOR FILING DATE: 2003-01-07
PRIOR APPLICATION NUMBER: DE 10245779.4
PRIOR FILING DATE: 2002-10-01
PRIOR FILING DATE: 2002-10-01
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APPLICANT: FORKENS, John
APPLICANT: HARBECK, Nadia
                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 2147
SEQ ID NO 307
                                                                                                Query Match 4.0%;
Best Local Similarity 43.7%;
Matches 178; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                   OTHER INFORMATION: chemically treated genomic DNA-10-517-441-307
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APPLICANT: HOEFLER, Heinz
TITLE OF INVENTION: Method and nucleic acids for the improved treatment of breast cel
TITLE OF INVENTION: proliferative disorders
FILE REFERENCE: 47675-93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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CURRENT FILING DATE: 2004-12-11
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                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                   LENGTH: 8093
                               1467 ATTGTATAATATTTGAGAGGATTTATGAATATAAATTGGGTTTTGTTTATAATTAGTTTT
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272 CTGCACTATTAGGTGAAAATGGTGATCAAATTTCTAAAGATATGGCAAATGGTTTTTATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  307, Application US/10517441
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MODEL, Fabian
NIMMRICH, Inko
RUJAN, Tamas
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SCHMITT, Manfred
LOOK, Maxime P.
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MAIER, Sabine
                                                                                                  <u>,</u>
                                                                                              Score 40.6; DB 7; Length 8093; Pred. No. 2:7; 0; Mismatches 229; Indels 0
                                                                                                                                                                                  (Homo sapiens)
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                                                                                                Gaps
                               1526
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                                                                                                                 Matches 178;
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; FEATURE:
; OTHER INFORMATION: chemically treated
US-10-517-441-441
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US-10-517-441-441
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SEQ ID NO 441
LENGTH: 12610
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TITLE OF INVENTION: Method and nucleic acids for the improved treatment of breast TITLE OF INVENTION: Details retive disorders
FILE OF INVENTION: Details retive disorders
FILE REFERENCE: 47675-93
CURRENT FILING DATE: 2004-12-11
PRIOR APPLICATION NUMBER: US/10/517,441
CURRENT FILING DATE: 2003-10-01
PRIOR FILING DATE: 2003-04-17
PRIOR APPLICATION NUMBER: DE 10317955.0
PRIOR APPLICATION NUMBER: DE 1031096.8
PRIOR FILING DATE: 2003-01-07
PRIOR APPLICATION NUMBER: DE 10245779.4
PRIOR FILING DATE: 2003-01-07
PRIOR FILING DATE: 2003-01-01
PRIOR FILING DATE: 2003-01-01
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                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                    Local Similarity
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291 TGGTGATCAAAGTTTCTAAAGATATGGCAAATGGTTTTTATAAGAATGAACTGGATTTTGA 350
                                                                                                                231 TGCTGAGAGAATTTATCAAGAATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAA 290
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                                                                     AAGTGTTTATTTTAGGTÄGGATTTTTTAAAATAAAGTAGTAATATTTACGTAGATAAGAT 1826
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NIMMRICH, Inko
RUJAN, Tamas
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SCHMITT, Manfred
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                                                                                                                                                              Conservative
                                                                                                                                                                                 4.0%;
                                                                                                                                                              0,
                                                                                                                                                                                 Score 40.4; D
Pred. No. 3.2;
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US-10-517-441-715
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                                                                                                                                                             Query Match 4.0%;
Best Local Similarity 45.9%;
                                                                                                                                                                                                                                                                                                                            SEQ ID NO 715
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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CURRENT FILING DATE: 2004-12-11
PRIOR APPLICATION NUMBER: PCT/EP2003/010881
PRIOR FILING DATE: 2003-10-01
PRIOR APPLICATION NUMBER: DE 10317955.0
PRIOR FILING DATE: 2003-04-17
PRIOR FILING DATE: 2003-01-07
PRIOR FILING DATE: 2003-01-07
PRIOR FILING DATE: 2003-01-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: LOOK, Maxime P.
APPLICANT: MARX, Almuth
APPLICANT: MOFFLER, Heinz
TITLE OF INVENTION: Method and nucleic acids for the improved treatment of breast cel
TITLE OF INVENTION: proliferative disorders
FILE REFERENCE: 47675-93
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NUMBER OF SEQ ID NOS: 2147
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                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: DE 10245779.4
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                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                             LENGTH: 12610
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                                                                                                          231 TGCTGAGAGAATTTATCAAGAATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAA
                                                                         MARTENS, John
MODEL, Fabian
NIMMRICH, Inko
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SCHMITT, Armin
SCHMITT, Manfred
                                   TGGTGATCAAATTTCTAAAGATATGGCAAATGGTTTTTATAAGAATGAACTGGATTTTGA
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MAIER, Sabine
                                                                                                                                                Conservative
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                                                                                                                                            Score 40.4; DB 7;
Pred. No. 3.2;
0; Mismatches 206;
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APPLICANT: ASHIKARI, TOSHIHIKO
TITLE OF INVENTION: METHODS FOR ANALYZING GENES OF INDUSTRIAL YEASTS
FILE REFERENCE: 030685-043
CURRENT APPLICATION NUMBER: US/10/932,182A
CURRENT FILING DATE: 2004-09-02
NUMBER OF SEQ ID NOS: 197023
SOFTWARE: Patentin version 3.3
SOFTWARE: Patentin version 3.3
SOFTWARE: Balo
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 243; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Saccharomyces pastorianus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          70 ATAGTACGTATACTAAACTTTAAGCAAACAGATGAGGGTCATACAGCATCATATGGAATT 129
                                                                                                                                                                                                                              GAATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAAATTTTCTAAA 309
TTATTAGCACTATATTTTTTCTCTGTACAGGAAATTCCACTGGAGGAAAATCAACAATCA
                                                    GATATGGCAAATGGTTTTTATAAGAATGAACTGGATTTTGAAGGTCAATATCCTCAAAAC
                                                                                                                                                                                                  ATTTATAGGGCTTACCTTGTTTAACTAATAATACGTAAATATTTAACTTAGCTCTCAAAC
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                                                                                        ATTTGGAATGTTCCTGAGCTTGAAAATAAACCATTGAGTGCTTATTCAGATGACGATAAA
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Pred. No. 2.2;
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US-10-868-397-1
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                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Sim
Matches 199;
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SEQ ID NO 1
LENGTH: 1691140
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PRIOR
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TITLE OF INVENTION: SUSCEPTIBILITY GENE FOR
TITLE OF INVENTION: METHOD OF TREATMENT
FILE REFERENCE: 2345.2010-012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Gretarsdottir, Solveig
APPLICANT: Thorleifsson, Gudmar
APPLICANT: Gilcher, Jeffrey R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 2003-08-27
PRIOR APPLICATION NUMBER: 10/419,723
PRIOR FILING DATE: 2003-04-18
PRIOR APPLICATION NUMBER: 10/255,120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/868,397
CURRENT FILING DATE: 2004-06-14
PRIOR APPLICATION NUMBER: PCT/US03/29906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2002-09-25
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PRIOR APPLICATION NUMBER: 10/650,120
                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
LOCATION: 1-1691140
                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 10/067,514 FILING DATE: 2002-02-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 09/811,352
                    383
                                                                                                                                                                                                                                                                                  143 AGAAAATAATATTAGCTTACGCTTTGGCTGTAAGTGGTATTCATAATGTATCTAAACTTC
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                                                                                                                                                                                                             CTGATGACTATTATAAGAATAAAGAGACTGCTGAGAGAATTTATCAAGAATATATGTCTA
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                                                  GTATTTGAAAAGCTCAACCTCAGCTTCCCCCTGTTGTTTATCTAGAATATTTAAGACTTT
                                                                                                                 AGAAAGCCTAAATAGTATTTTAAAAATAAAGCTTCCATATATTCAATCTTATTGGAAAAA
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                                                                                  GTTTTTATAAGAATGAACTGGATTTTGAAGGTCAATATCCTCAAAACATTTGGAATGTTC
                                                                                                                                                ATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAAATTTCTAAAGATATGGCAAATG
                                                                                                                                                                                 CCTGGACATTGAGTAACAAATAAGTTAATAAAGCAATAGTTTGTTATTTGATTTTATATC
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Pred. No. 9.1;
0; Mismatches 265; Indels 0;
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US-10-932-182A-76518
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US-11-021-837-43
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Matches
                                                                                 Sequence 76518, Application US/10932182A GENERAL INFORMATION:
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APPLICANT: NAKAO, YOSHIHIRO
APPLICANT: NAKAMURA, NORII
APPLICANT: KODAMA, YUKIKO
APPLICANT: FUJIMURA, TOMOD
APPLICANT: ASHIKARI, TOSHI
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                                                                   YOSHIHIRO
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; TYPE: DNA ; ORGANISM: Staphylococcus saprophyticus US-11-021-837-43
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 43
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APPLICANT: Manning, Paul
APPLICANT: Mclaughlin, Robert
APPLICANT: Mclaughlin, Robert
APPLICANT: MacCormack, Kathleen
TITLE OF INVENTION: Staphlococcus Saprophyticus Nucleic Acids and Polypeptides
FILE REFERENCE: 101324-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/11/021,837 CURRENT FILING DATE: 2004-12-23 PRIOR APPLICATION NUMBER: US 60/533534 PRIOR FILING DATE: 2003-12-31 PRIOR APPLICATION NUMBER: US 60/600680 PRIOR APPLICATION NUMBER: US 60/600680 PRIOR FILING DATE: 2003-08-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
mes 116; Conserv
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14253 AAA 14255
                                                                                                                                                                                                                                                                                                                                                                   14073 AGATGGTAGTAGGTATTAAGTATTATATTAĞTTAATACTGCCACAAAATGTGGCTTAAGT 14132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    67 GATATAGTACGTATACTAAACTTTAAGCAAACAGATGAGGGTCATACAGCATCATATGGA
                                                                                                                                                                                                                                          GGTCAATTTGAAGAATTAGAAGAATATTACCAAAAATATAAAGAGCAAGGGCTAATTATA 14192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GTTATAAAAGGAGTGCTATAAAATAACTAGTAATATTTATGATATACAAGTTGTTAACAA 14072
                                                                                                                        CTTGGATTTCCTTGTAATCAATTTGCTAAACAAAAACCAGGTACTGATGACCAAATAGCT 14252
                                                                                                                                                                              CAAGAATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAAATTTCT
                                                                                                                                                                                                                                                                                          AATGTATCTAAACTTCCTGATGACTATTATAAGAATAAAGAGACTGCTGAGAGAATTTAT
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US-10-517-441-414/c
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; ORGANISM: Saccharomyces pastorianus
US-10-932-182A-76518
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CURRENT APPLICATION NUMBER: US/10/932,182A
CURRENT FILING DATE: 2004-09-02
RUMBER OF SEQ ID NOS: 197023
SOFTWARE: PATENTIN VERSION 3.3
SQI DNO 76518
LENGTH: 2460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 414, Application US/10517441
GENERAL INFORMATION:
APPLICANT: FOEKENS, John
APPLICANT: HARBECK, Nadia
APPLICANT: KOENIG, Thomas
                                                                                                                                                  SEQ ID NO 414
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Best Local Similarity
   Query Match
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TITLE OF INVENTION: Method and nucleic acids
TITLE OF INVENTION: proliferative disorders
FILE REFERENCE: 47675-93
CURRENT APPLICATION NUMBER: US/10/517,441
CURRENT FILING DATE: 2004-12-11
                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: PCT/EP2003/010881
PRIOR FILING DATE: 2003-10-01
PRIOR APPLICATION NUMBER: DE 10317955.0
PRIOR FILING DATE: 2003-04-17
PRIOR APPLICATION NUMBER: DE 10300096.8
PRIOR FILING DATE: 2003-01-07
                                                                                                                                                                 PRIOR APPLICATION NUMBER: DE 10245779.4
PRIOR FILING DATE: 2002-10-01
NUMBER OF SEQ ID NOS: 2147
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                                                                                    ORGANISM: Artificial Sequence
                                                       OTHER INFORMATION: chemically treated genomic DNA
                                                                             FEATURE:
                                                                                                                 TYPE: DNA
                                                                                                                                 LENGTH: 19634
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TER, Sabine
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SCHMITT, Manfred
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Pred. No. 3.7;
0; Mismatches
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 Score 39.2;
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                                                                                                                                                                                                                                                                                                                                                                                                                  the improved treatment
Length 19634;
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                                                       (Homo sapiens)
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                                                                                                                                                                                                                                                                                                                                                                                                                  of breast
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TITLE OF INVENTION: proliferative disorders FILE REFERENCE: 47675-93
CURRENT APPLICATION NUMBER: US/10/517,441
CURRENT FILING DATE: 2004-12-11
PRIOR APPLICATION NUMBER: PCT/EP2003/01081
PRIOR FILING DATE: 2003-10-01
PRIOR PRICINGION NUMBER: DE 10317955.0
PRIOR APPLICATION NUMBER: DE 10317955.0
PRIOR FILING DATE: 2003-04-17
PRIOR APPLICATION NUMBER: DE 1030096.8
PRIOR APPLICATION NUMBER: DE 1030096.8
PRIOR FILING DATE: 2003-01-07
PRIOR APPLICATION NUMBER: DE 10345779.4
PRIOR FILING DATE: 2002-10-01
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SEQ ID NO 688
LENGTH: 19634
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                                                                                        Query Match 3.9%;
Best Local Similarity 44.7%;
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                                                                       Matches 152;
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TITLE OF INVENTION: Method and nucleic acids for the improved treatment of breast celette of properties. Proliferative disorders
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                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                OTHER INFORMATION: chemically treated genomic DNA
                                                                                                                                                                                                          FEATURE:
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245 ATCAAGAATATATGTCTAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAAATTT 304
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SCHMITT, Manfred
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NIMMRICH, Inko
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                                                                       Conservative
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                                                                     Score 39.2; DB 7;
Pred. No. 6.5;
0; Mismatches 188;
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                                                                       Indels
                                                                                                                                                                                  (Homo sapiens)
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; FEATURE:
; NAME/KEY: midc feature
; LOCATION: (1)...(397658)
; OTHER INFORMATION: n = A,T,C
US-11-044-879-3
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US-11-044-879-3/d
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GENERAL INFORMATION:
APPLICANT: ZHANG, HONDYU et al.
APPLICANT: ZHANG, HONDYU et al.
TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: CL001172CON
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
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Best Local
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CURRENT FILING DATE: 2005-01-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
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340738 TÍCTAATTTAACGTGTAATAGAGATAAGATCCAAGTGGTGAGACAGAACAACAGGAGCAG 340679
                                                                                                                                  340858
                                                                                                                                                                                              341038 TATGTATGTTACATATACACATACTATGTATGTATGTATTATATATTCAGCACATATAATAT
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                              ATATTTTTTCTC-TGTACAGGAAATTCCACTGGAGGAAAATCAACAATCAAATGCCGCAA
                                                                CATAGTCCCTGTGTTGAAGAAGATGACTCTAACTGAGGAGGTAAATAACTAATTGCCTAA 340739
                                                                                                                             TAATCTTTCATCTGCACTATTAGGTGAAAATGGTGATCAAATTTCTAAAGATATGGCAAA
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llarity 45.5%;
Conservative
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Pred. No. 11;
0; Mismatches 253;
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Sequence 1254, Application US/10450763

GENERAL INFORMATION:
APPLICANT: Hyseq, Inc
TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POL
FILE REFERENCE: 790CIP3/US
CURRENT FILING DATE: 2003-06-11
PRIOR APPLICATION NUMBER: US/10/450,763
CURRENT FILING DATE: 2001-06-11
PRIOR APPLICATION NUMBER: PCT/US01/08631
PRIOR FILING DATE: 2001-03-30
PRIOR PILING DATE: 2001-03-31
PRIOR APPLICATION NUMBER: 09/540,217
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR PILING DATE: 2000-08-23
PRIOR FILING DATE: 2000-08-23
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Search completed: March Job time: 280 secs
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US-10-450-763-12549
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SEQ ID NO 12549
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (916)..(215)
OTHER INFORMATION: 57% homologous to Plasmodium lophurae histidine-rich
OTHER INFORMATION: protein, accession number X01469, Smith-Waterman Score=915.
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NAME/KEY: SIMILAR
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                                                                                AGTGATGGTAATGATGGTGATGATGATA
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                   2005,
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